

Serial Number: 09/485,951

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☒ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☒ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: 10
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/485,951

DATE: 01/30/2001
TIME: 10:22:53

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FEB - 1 2001

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Output Set: N:\CRF3\01302001\I485951.raw

TECH CENTER 1600/2900

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3 <110> APPLICANT: Seisi Kato
4   Yamaguchi Kimura
5   Shingo Sekine
6   Kouju Kamata
8 <120> TITLE OF INVENTION: HUMAN GALECTIC-9-LIKE PROTEINS AND cDNA ENCODING THESE
9   PROTEINS
11 <130> FILE REFERENCE: GIN-6707CPUS
13 <140> CURRENT APPLICATION NUMBER: 09/485,951
14 <141> CURRENT FILING DATE: 2000-02-17
16 <150> PRIOR APPLICATION NUMBER: 9-226468
17 <151> PRIOR FILING DATE: 1997-08-22
19 <150> PRIOR APPLICATION NUMBER: PCT/JP98/03670
20 <151> PRIOR FILING DATE: 1998-08-19
22 <160> NUMBER OF SEQ ID NOS: 11
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54 Val Asn Gly Thr Val Leu Ser Ser Ser Gly Thr Arg Phe Ala Val Asn
55      35      40      45
57 Phe Gln Thr Gly Phe Ser Gly Asn Asp Ile Ala Phe His Phe Asn Pro
58      50      55      60
60 Arg Phe Glu Asp Gly Gly Tyr Val Val Cys Asn Thr Arg Gln Asn Gly
61 65      70      75      80
63 Ser Trp Gly Pro Glu Arg Lys Thr His Met Pro Phe Gln Lys Gly
64      85      90      95
66 Met Pro Phe Asp Leu Cys Phe Leu Val Gln Ser Ser Asp Phe Lys Val
67      100     105     110
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72 His Arg Val Asp Thr Ile Ser Val Asn Gly Ser Val Gln Leu Ser Tyr
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76 145 150 155 160
78 Thr Val Pro Phe Ser Gln Pro Val Cys Phe Pro Pro Arg Pro Arg Gly
79 165 170 175
81 Arg Arg Gln Lys Pro Pro Gly Val Trp Pro Ala Asn Pro Ala Pro Ile
82 180 185 190
84 Thr Gln Thr Val Ile His Thr Val Gln Ser Ala Pro Gly Gln Met Phe
85 195 200 205
87 Ser Thr Pro Ala Ile Pro Pro Met Met Tyr Pro His Pro Ala Tyr Pro
88 210 215 220
90 Met Pro Phe Ile Thr Thr Ile Leu Gly Gly Leu Tyr Pro Ser Lys Ser
91 225 230 235 240
93 Ile Leu Leu Ser Gly Thr Val Leu Pro Ser Ala Gln Arg Phe His Ile
94 245 250 255
96 Asn Leu Cys Ser Gly Asn His Ile Ala Phe His Leu Asn Pro Arg Phe
97 260 265 270
99 Asp Glu Asn Ala Val Val Arg Asn Thr Gln Ile Asp Asn Ser Trp Gly
100 275 280 285
102 Ser Glu Glu Arg Ser Leu Pro Arg Lys Met Pro Phe Val Arg Gly Gln
103 290 295 300
105 Ser Phe Ser Val Trp Ile Leu Cys Glu Ala His Cys Leu Lys Val Ala
106 305 310 315 320
108 Val Asp Gly Gln His Leu Phe Glu Tyr Tyr His Arg Leu Arg Asn Leu
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139 agtggaaacca ggtttgctgt gaactttcag actggcttca gtggaaatga cattgccttc 180
141 cacttcaacc ctcggttga agatggaggg tacgtggtgt gcaacacgag gcagaacgga 240
143 agctgggggc ccaggagag gaagacacac atgcctttcc agaaggggat gccctttgac 300
145 ctctgcttcc tggtcagag ctacagatttc aaggatgatg tgaacgggat cctcttcgtg 360
147 cagtacttcc accgcgtgcc ctccaccgt gtggacacca tctccgtcaa tggtctgtg 420
149 cagctgtcct acatcagctt ccagaacccc cgcacagtc ctgttcagcc tgccttctcc 480
151 acggtgccgt tctccagcc tgtctgttcc ccaccaggc ccagggggcg cagacaaaaa 540

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DATE: 01/30/2001

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Output Set: N:\CRF3\01302001\I485951.raw

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157 cccgcctatc cgatgccttt catcaccacc attctgggag ggctgtaccc atccaagtcc 720
159 atcctcctgt caggcactgt cctgccaggt gctcagaggt tccacatcaa cctgtgctct 780
161 gggaaccaca tcgccttcca cctgaacccc cgttttgatg agaatgctgt ggtccgcaac 840
163 acccagatcg acaactcctg ggggtctgag gacggaagtc tgccccgaaa aatgcccttc 900
165 gtccgtggcc agagcttctc agtgtggatc ttgtgtgaag ctcaactgct caaggtggcc 960
167 gtggatggtc agcacctgtt tgaatactac catcgccctg ggaacctgcc caccatcaac 1020
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184 ggccacagag gcggcggaga g atg gcc ttc agc ggt tcc cag gct ccc tac 111
185 Met Ala Phe Ser Gly Ser Gln Ala Pro Tyr
186 1 5 10
188 ctg agt cca gct gtc ccc ttt tct ggg act att caa gga ggt ctc cag 159
189 Leu Ser Pro Ala Val Pro Phe Ser Gly Thr Ile Gln Gly Gly Leu Gln
190 15 20 25
192 gac gga ctt cag atc act gtc aat ggg acc gtt ctc agc tcc agt gga 207
193 Asp Gly Leu Gln Ile Thr Val Asn Gly Thr Val Leu Ser Ser Ser Gly
194 30 35 40
196 acc agg ttt gct gtg aac ttt cag act ggc ttc agt gga aat gac att 255
197 Thr Arg Phe Ala Val Asn Phe Gln Thr Gly Phe Ser Gly Asn Asp Ile
198 45 50 55
200 gcc ttc cac ttc aac cct cgg ttt gaa gat gga ggg tac gtg gtg tgc 303
201 Ala Phe His Phe Asn Pro Arg Phe Glu Asp Gly Gly Tyr Val Val Cys
202 60 65 70
204 aac acg agg cag aac gga agc tgg ggg ccc gag gag agg aag aca cac 351
205 Asn Thr Arg Gln Asn Gly Ser Trp Gly Pro Glu Glu Arg Lys Thr His
206 75 80 85 90
208 atg cct ttc cag aag ggg atg ccc ttt gac ctc tgc ttc ctg gtg cag 399
209 Met Pro Phe Gln Lys Gly Met Pro Phe Asp Leu Cys Phe Leu Val Gln
210 95 100 105
212 agc tca gat ttc aag gtg atg gtg aac ggg atc ctc ttc gtg cag tac 447
213 Ser Ser Asp Phe Lys Val Met Val Asn Gly Ile Leu Phe Val Gln Tyr
214 110 115 120
216 ttc cac cgc gtg ccc ttc cac cgt gtg gac acc atc tcc gtc aat ggc 495
217 Phe His Arg Val Pro Phe His Arg Val Asp Thr Ile Ser Val Asn Gly
218 125 130 135
220 tct gtg cag ctg tcc tac atc agc ttc cag aac ccc cgc aca gtc cct 543
221 Ser Val Gln Leu Ser Tyr Ile Ser Phe Gln Asn Pro Arg Thr Val Pro
222 140 145 150
224 gtt cag cct gcc ttc tcc acg gtg ccg ttc tcc cag cct gtc tgt ttc 591

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/485,951

DATE: 01/30/2001

TIME: 10:22:53

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Output Set: N:\CRF3\01302001\I485951.raw

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229 Pro Pro Arg Pro Arg Gly Arg Arg Gln Lys Pro Pro Gly Val Trp Pro
230                               175                               180                               185
232 gcc aac ccg gct ccc att acc cag aca gtc atc cac aca gtg cag agc 687
233 Ala Asn Pro Ala Pro Ile Thr Gln Thr Val Ile His Thr Val Gln Ser
234                               190                               195                               200
236 gcc cct gga cag atg ttc tct act ccc gcc atc cca cct atg atg tac 735
237 Ala Pro Gly Gln Met Phe Ser Thr Pro Ala Ile Pro Pro Met Met Tyr
238                               205                               210                               215
240 ccc cac ccc gcc tat ccg atg cct ttc atc acc acc att ctg gga ggg 783
241 Pro His Pro Ala Tyr Pro Met Pro Phe Ile Thr Thr Ile Leu Gly Gly
242                               220                               225                               230
244 ctg tac cca tcc aag tcc atc ctc ctg tca ggc act gtc ctg ccc agt 831
245 Leu Tyr Pro Ser Lys Ser Ile Leu Leu Ser Gly Thr Val Leu Pro Ser
246 235                               240                               245                               250
248 gct cag agg ttc cac atc aac ctg tgc tct ggg aac cac atc gcc ttc 879
249 Ala Gln Arg Phe His Ile Asn Leu Cys Ser Gly Asn His Ile Ala Phe
250                               255                               260                               265
252 cac ctg aac ccc cgt ttt gat gag aat gct gtg gtc cgc aac acc cag 927
253 His Leu Asn Pro Arg Phe Asp Glu Asn Ala Val Val Arg Asn Thr Gln
254                               270                               275                               280
256 atc gac aac tcc tgg ggg tct gag gag cga agt ctg ccc cga aaa atg 975
257 Ile Asp Asn Ser Trp Gly Ser Glu Glu Arg Ser Leu Pro Arg Lys Met
258                               285                               290                               295
260 ccc ttc gtc cgt ggc cag agc ttc tca gtg tgg atc ttg tgt gaa gct 1023
261 Pro Phe Val Arg Gly Gln Ser Phe Ser Val Trp Ile Leu Cys Glu Ala
262                               300                               305                               310
264 cac tgc ctc aag gtg gcc gtg gat ggt cag cac ctg ttt gaa tac tac 1071
265 His Cys Leu Lys Val Ala Val Asp Gly Gln His Leu Phe Glu Tyr Tyr
266 315                               320                               325                               330
268 cat cgc ctg agg aac ctg ccc acc atc aac aga ctg gaa gtg ggg ggc 1119
269 His Arg Leu Arg Asn Leu Pro Thr Ile Asn Arg Leu Glu Val Gly Gly
270                               335                               340                               345
272 gac atc cag ctg acc cat gtg cag aca taggcggcctt cctggccctg 1166
273 Asp Ile Gln Leu Thr His Val Gln Thr
274                               350                               355
276 gggccggggg ctgggggtgtg gggcagtcctg ggctcctctca tcatcccccac ttcccaggcc 1226
278 cagcctttcc aaccctgcct gggatctggg ctttaatgca gaggccatgt ccttgtcttg 1286
280 tcctgcttct ggctacagcc accctggaac ggagaaggca gctgacgggg attgccttcc 1346
282 tcagccgcag cagcacctgg ggctccagct gctggaatcc taccatccca ggaggcaggc 1406
284 acagccaggg agaggggagg agtgggcagt gaagatgaag ccccatgctc agtcccctcc 1466
286 catcccccac gcagctccac cccagtccca agccaccagc tgtctgctcc tgggtgggagg 1526
288 tggcctctc agccctcct ctctgacctt taacctcact ctcacctgc accgtgcacc 1586
290 aacccttcac cctccttgga aagcaggcct gatggcttcc cactggcctc caccacctga 1646
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294 aaatgcttgt tggcacatt
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RAW SEQUENCE LISTING

DATE: 01/30/2001

PATENT APPLICATION: US/09/485,951

TIME: 10:22:53

Input Set : A:\Pto.amc

Output Set: N:\CRF3\01302001\I485951.raw

298 <211> LENGTH: 355

299 <212> TYPE: PRT

300 <213> ORGANISM: Homo sapiens

302 <400> SEQUENCE: 6

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307           20           25           30
309 Val Asn Gly Thr Val Leu Ser Ser Ser Gly Thr Arg Phe Ala Val Asn
310           35           40           45
312 Phe Gln Thr Gly Phe Ser Gly Asn Asp Ile Ala Phe His Phe Asn Pro
313           50           55           60
315 Arg Phe Glu Asp Gly Gly Tyr Val Val Cys Asn Thr Arg Gln Asn Gly
316  65           70           75           80
318 Ser Trp Gly Pro Glu Glu Arg Lys Thr His Met Pro Phe Gln Lys Gly
319           85           90           95
321 Met Pro Phe Asp Leu Cys Phe Leu Val Gln Ser Ser Asp Phe Lys Val
322           100          105          110
324 Met Val Asn Gly Ile Leu Phe Val Gln Tyr Phe His Arg Val Pro Phe
325           115          120          125
327 His Arg Val Asp Thr Ile Ser Val Asn Gly Ser Val Gln Leu Ser Tyr
328           130          135          140
330 Ile Ser Phe Gln Asn Pro Arg Thr Val Pro Val Gln Pro Ala Phe Ser
331 145           150           155           160
333 Thr Val Pro Phe Ser Gln Pro Val Cys Phe Pro Pro Arg Pro Arg Gly
334           165          170          175
336 Arg Arg Gln Lys Pro Pro Gly Val Trp Pro Ala Asn Pro Ala Pro Ile
337           180          185          190
339 Thr Gln Thr Val Ile His Thr Val Gln Ser Ala Pro Gly Gln Met Phe
340           195          200          205
342 Ser Thr Pro Ala Ile Pro Pro Met Met Tyr Pro His Pro Ala Tyr Pro
343           210          215          220
345 Met Pro Phe Ile Thr Thr Ile Leu Gly Gly Leu Tyr Pro Ser Lys Ser
346 225           230           235           240
348 Ile Leu Leu Ser Gly Thr Val Leu Pro Ser Ala Gln Arg Phe His Ile
349           245          250          255
351 Asn Leu Cys Ser Gly Asn His Ile Ala Phe His Leu Asn Pro Arg Phe
352           260          265          270
354 Asp Glu Asn Ala Val Val Arg Asn Thr Gln Ile Asp Asn Ser Trp Gly
355           275          280          285
357 Ser Glu Glu Arg Ser Leu Pro Arg Lys Met Pro Phe Val Arg Gly Gln
358           290          295          300
360 Ser Phe Ser Val Trp Ile Leu Cys Glu Ala His Cys Leu Lys Val Ala
361 305           310           315           320
363 Val Asp Gly Gln His Leu Phe Glu Tyr Tyr His Arg Leu Arg Asn Leu
364           325          330          335
366 Pro Thr Ile Asn Arg Leu Glu Val Gly Gly Asp Ile Gln Leu Thr His
367           340          345          350
369 Val Gln Thr

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VERIFICATION SUMMARY DATE: 01/30/2001
PATENT APPLICATION: US/09/485,951 TIME: 10:22:54

Input Set : A:\Pto.amc
Output Set: N:\CRF3\01302001\I485951.raw

1642

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/485,951 DATE: 01/23/2001
 TIME: 14:39:46

Input Set : A:\seqlist.txt
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**Does Not Comply
 Corrected Diskette Needed**

3 <110> APPLICANT: Seisi Kato
 4 Yamaguchi Kimura
 5 Shingo Sekine
 6 Kouju Kamata
 8 <120> TITLE OF INVENTION: HUMAN GALECTIC-9-LIKE PROTEINS AND CDNA ENCODING THESE
 9 PROTEINS
 11 <130> FILE REFERENCE: GIN-6707CPUS
 13 <140> CURRENT APPLICATION NUMBER: 09/485,951
 14 <141> CURRENT FILING DATE: 2000-02-17
 16 <150> PRIOR APPLICATION NUMBER: 9-226468
 17 <151> PRIOR FILING DATE: 1997-08-22
 19 <150> PRIOR APPLICATION NUMBER: PCT/JP98/03670
 20 <151> PRIOR FILING DATE: 1998-08-19
 22 <160> NUMBER OF SEQ ID NOS: 11
 24 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

E--> 531 <210> SEQ ID NO: 11
 532 <211> LENGTH: 30
 533 <212> TYPE: DNA
 534 <213> ORGANISM: Artificial Sequence
 536 <220> FEATURE:
 537 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/485,951

DATE: 01/23/2001

TIME: 14:39:47

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L:543 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO:11